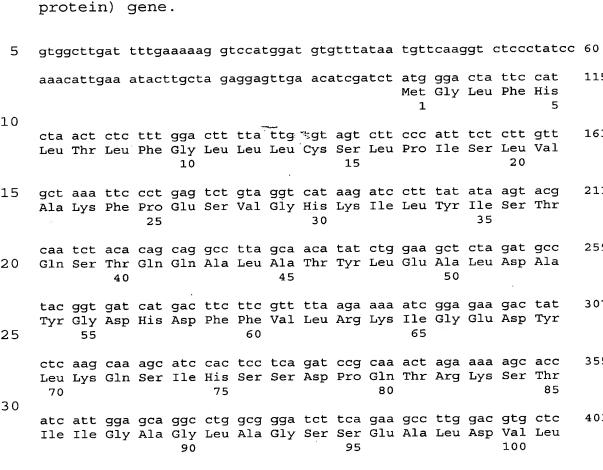
Figure 1. Sequence of C. pneumoniae OMP (outer membrane



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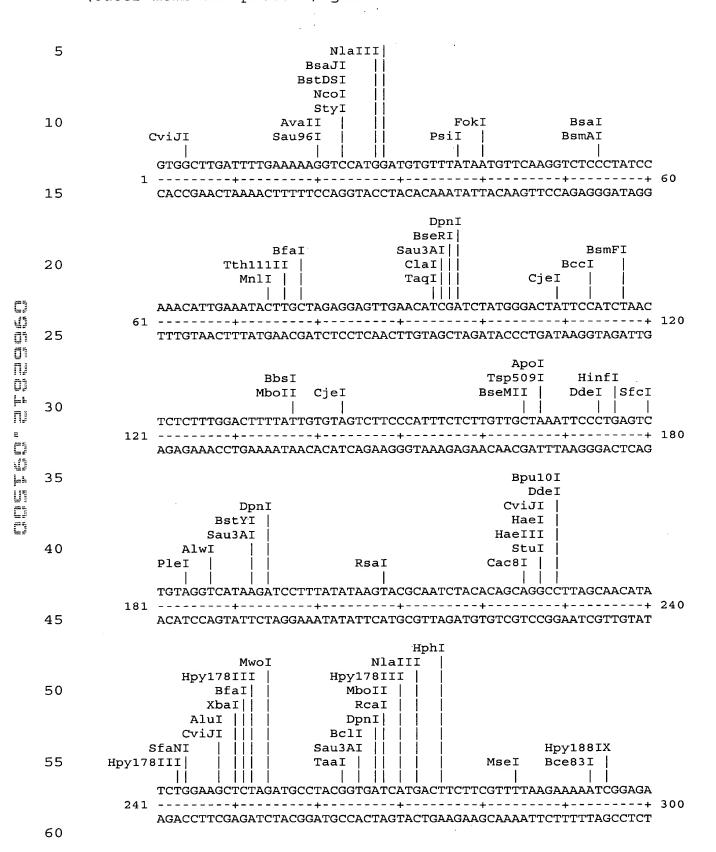
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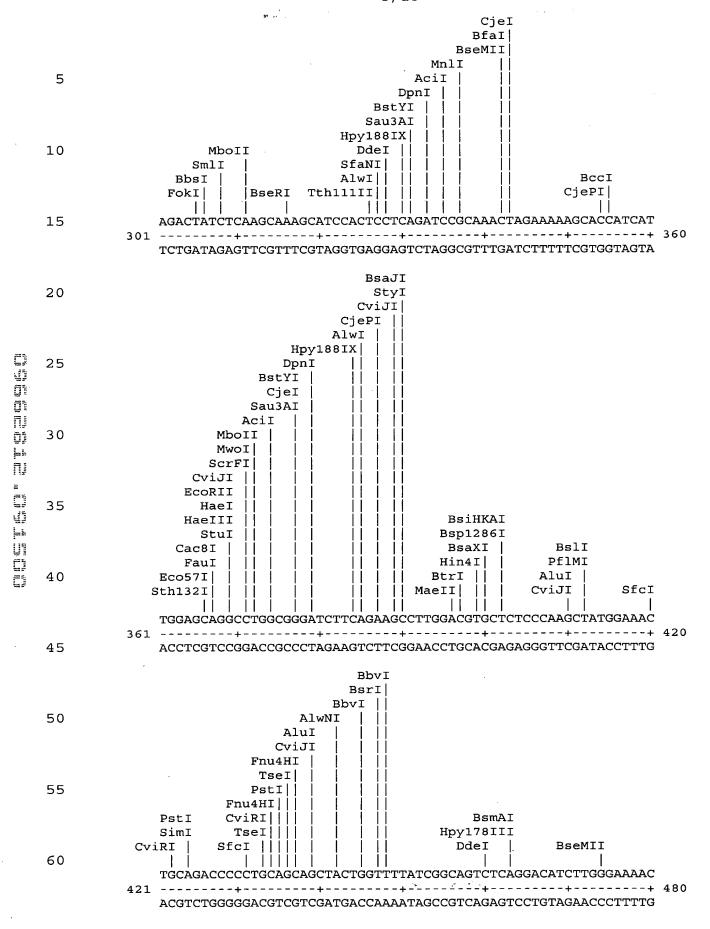
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5	acg Thr 230	agt Ser	gcg Ala	tct Ser	cct Pro	caa Gln 235	gat Asp	caa Gln	gaa Glu	gct Ala	att Ile 240	ctt Leu	tat Tyr	gct Ala	tta Leu	999 Gly 245	835
10																	883
15	aag Lys	cct Pro	gat Asp	gtg Val 265	gat Asp	gtc Val	act Thr	tta Leu	gca Ala 270	gca Ala	gct Ala	caa Gln	gct Ala	tta Leu 275	att Ile	gct Ala	931
2.0	ttg Leu	gly ggg	aaa Lys 280	gaa Glu	gag Glu	gac Asp	gct Ala	ctt Leu 285	ccc Pro	gtg Val	ata Ile	aaa Lys	aag Lys 290	caa Gln	gca Ala	ctt Leu	979
20	gag Glu	gag Glu 295	cgg Arg	cct Pro	cga Arg	gcc Ala	ctg Leu 300	tat Tyr	gcc Ala	tta Leu	cgg Arg	cat His 305	cta Leu	ccc Pro	tct Ser	gag Glu	1027
25	ata Ile 310	gly ggg	att Ile	ccg Pro	att Ile	gcc Ala 315	ctg Leu	ccg Pro	ata Ile	ttc Phe	cta Leu 320	aaa Lys	act Thr	aag Lys	aac Asn	agc Ser 325	1075
30	gaa Glu	gcc Ala	aag Lys	ttg Leu	aat Asn 330	gta Val	gct Ala	tta Leu	gct Ala	ctc Leu 335	tta Leu	gag Glu	tta Leu	Gly aaa	tgt Cys 340	gac Asp	1123
35	acc Thr	cct Pro	aaa Lys	cta Leu 345	ctg Leu	gaa Glu	tac Tyr	att Ile	acc Thr 350	gaa Glu	agg Arg	ctt Leu	gtc Val	caa Gln 355	cca Pro	cat His	1171
40																	1219
40																	1267
45	gaa Glu 390	agg Arg	ttg Leu	ctc Leu	tcc Ser	aca Thr 395	acc Thr	cga Arg	ggt Gly	ctt Leu	gaa Glu 400	gag Glu	cag Gln	atc Ile	ctt Leu	acg Thr 405	1315
50																	1363
55	ctt Leu	ttg Leu	gcg Ala	agt Ser 425	cag Gln	aaa Lys	act Thr	cag Gln	ctt Leu 430	gcc Ala	act Thr	act Thr	gcg Ala	att Ile 435	tct Ser	ttt Phe	1411
	tta Leu	agt Ser	cac His 440	acc Thr	tca Ser	cat His	cag Gln	gaa Glu 445	gcc Ala	tta Leu	gat Asp	cta Leu	ctt Leu 450	ttc Phe	caa Gln	gct Ala	1459
00												Tyr	Ala				1507
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ggg att ccg Ile Gly Ile Pro 310 acc Cct aaa cta Lys Leu Ala Lys Leu 345 tat aat gag act Tyr Asn Glu Thr 360 40 aat tgg aag cgg Arg Arg 375 45 gaa agg ttg Ctc Glu Arg 275 45 gaa agg ttg ctc Glu Arg 390 ctt ttg ggg aag cgg Arg Ctc Glu Arg Leu Arg 390 ctt ttg gcg agt Ser Ser Ala Ser Arg Arg Arg Arg Arg Arg Arg Arg Arg Ar	Thr Ser Ala Ser Pro 230 10	Thr 230 Ser Ala Ser Pro Gln 235 10 Lys Leu Lys Asp Ggt Gggt Cag Gln 250 Gggt Ggg Gln 250 Ggg Ggg Gln 250 Ggg Ggg Gln 250 Ggg Ggg Ggg Ggg Gac Ggg Ggg Gac Gglu 260 Ggg Ggg Ggg Ggg Gac Gglu 280 Glu Asp 260 Ggg Ggg Ggg Ggg Gac Gglu 290 Ggg Ggg Ggg Ggg Gac Asp 290 Ggg Ggg Ggg Ggg Gac Asp 290 Ggg Asp 290 Ggg Ggg Ggg Gac Asp 290 Ggg	Thr 230 Ser Ala Ser Pro Gln Asp 235 aag ctt aag gat ggt gat ggt gat Lys Pro Asp 250 aag cct gat gtg gat ggt gat Cyal Thr 250 ttg ggg aaa gaa gaa gag gac gct Lys Glu Glu Glu Hys Glu	Thr 230 Ser Ala Ser Pro 235 Cla 235 Asp 235 Cla 235 Cla 235 Asp 235 Cla 235 Asp 235 Cla 235 Asp 240 Asp 240 <td> Thr Ser Ala Ser Pro Gln Asp Gln 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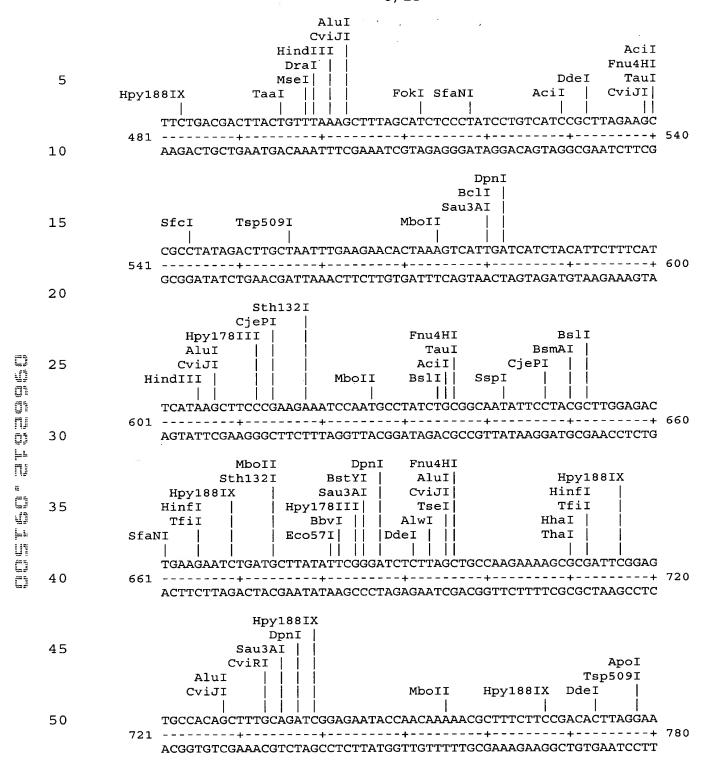
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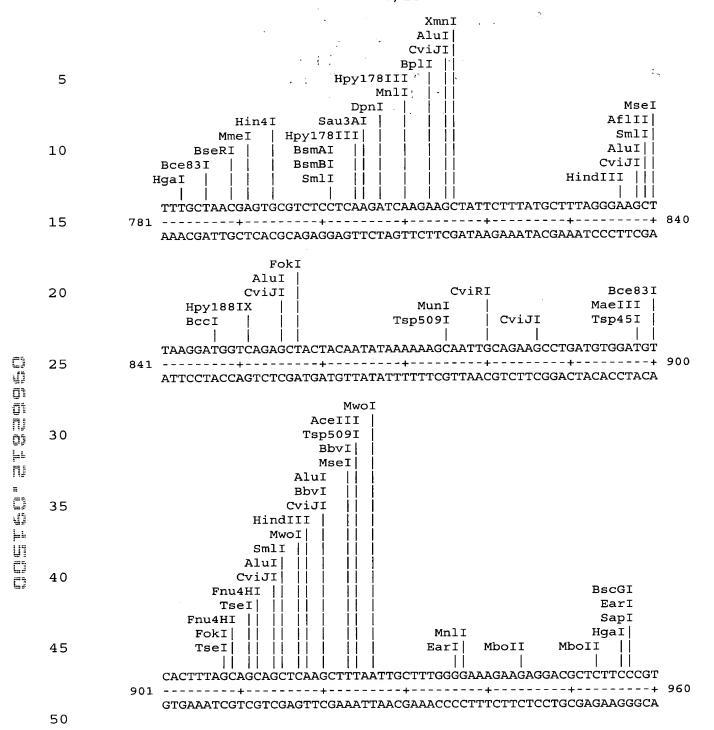
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5	tat Tyr	gca Ala	aaa Lys	aag Lys	cta Leu 490	att Ile	cag Gln	gaa Glu	acc Thr	ttg Leu 495	tta Leu	ttt Phe	gtg Val	gac Asp	acg Thr 500	gaa Glu	1603
10	aac Asn	caa Gln	aga Arg	ccc Pro 505	cat His	ccc Pro	agc Ser	atg Met	ccc Pro 510	tat Tyr	cta Leu	cgt [.] Arg	tat Tyr	cag Gln 515	gtc Val	acc Thr	1651
15	cca Pro	gaa Glu	agc Ser 520	cgt Arg	acg Thr	aag Lys	ctc Leu	atg Met 525	ttg Leu	gat Asp	att Ile	cta Leu	gag Glu 530	aca Thr	cta Leu	gcc Ala	1699
20	acc Thr	tcg Ser 535	aag Lys	tct Ser	tcc Ser	gaa Glu	gat Asp 540	atc Ile	cgt Arg	tta Leu	ttg Leu	ata Ile 545	caa Gln	ctg Leu	atg Met	acg Thr	1747
20	gaa Glu 550	gga Gly	gat Asp	gca Ala	aaa Lys	aat Asn 555	ttc Phe	cca Pro	gtc Val	ctt Leu	gca Ala 560	ggc Gly	tta Leu	ctc Leu	ata Ile	aaa Lys 565	1795
25		att gtg gag taaccccaac ctacgtctta tgaaacgttg cttcttattt Ile Val Glu															1844
	ctag	gctto	cct t	tgti	ctta	at g	ggtto	cctca	a gct	gate	gctt	tgad	ctcat	ca a	agag	gctgtg	1904
30	aaa																1907

Figure 2. Restriction enzyme analysis of the *C. pneumoniae* OMP (outer membrane protein) gene.

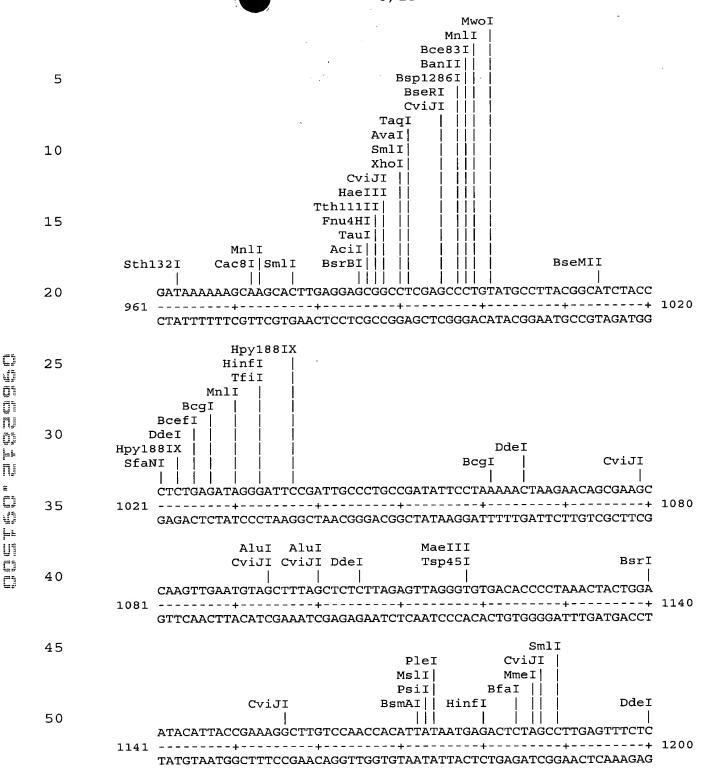


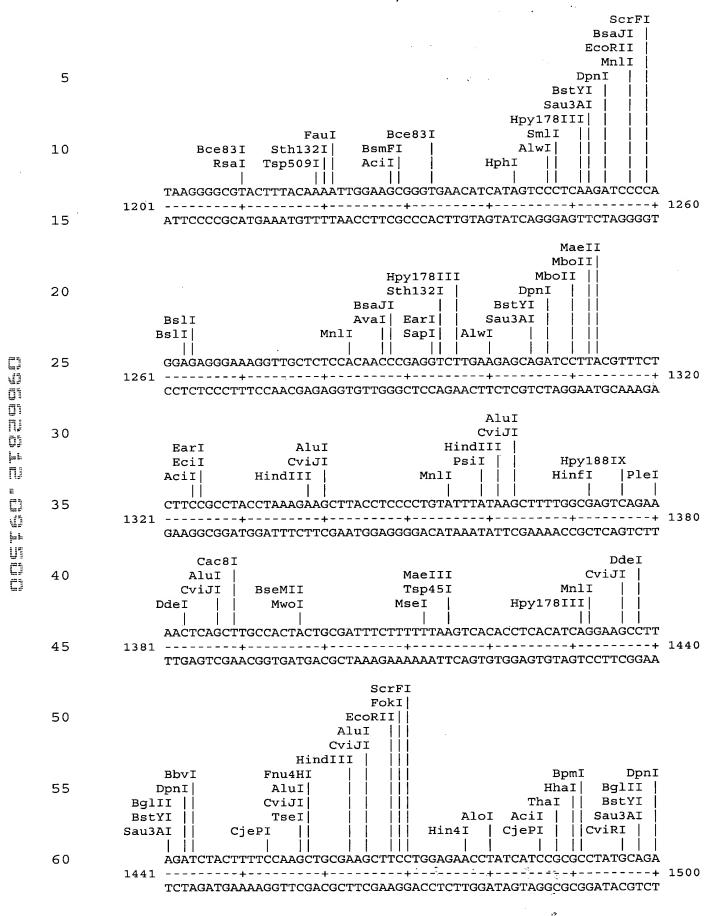




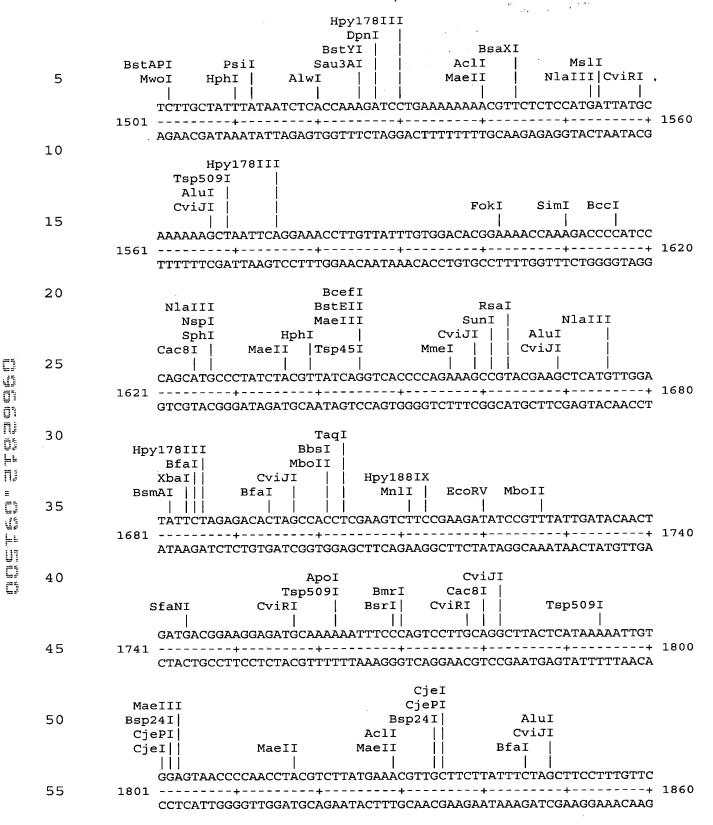


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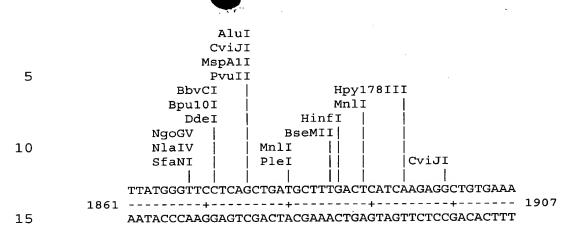


Figure 4: Protective efficacy of DNA immunization with pCAmg002.

